

GenCore version 4.5
Copyright (c) 1993 - 2000 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: August 14, 2002, 10:50:38 ; Search time 75.95 Seconds

(without alignments)
45.336 Million cell updates/sec

Title: US-09-785-059-2

Perfect score: 148

Sequence: 1 RYRVQACRAIRHIVRIRROGLRRIRRV 31

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 747574 seqs, 111073796 residues

Total number of hits satisfying chosen parameters: 747574

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :
A_Geneseq_032802:*

1: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1980.DAT:*
2: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1981.DAT:*
3: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1982.DAT:*
4: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1983.DAT:*
5: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1984.DAT:*
6: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1985.DAT:*
7: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1986.DAT:*
8: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1987.DAT:*
9: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1988.DAT:*
10: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1989.DAT:*
11: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1990.DAT:*
12: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1991.DAT:*
13: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1992.DAT:*
14: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1993.DAT:*
15: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1994.DAT:*
16: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1995.DAT:*
17: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1996.DAT:*
18: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1997.DAT:*
19: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1998.DAT:*
20: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA1999.DAT:*
21: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA2000.DAT:*
22: /SIDSI/gcgdata/hold-geneseq/geneeqp-emb1/AA2001.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	122	82.4	28	19	AAW47769
2	122	82.4	28	20	AAW47769
3	117	79.1	28	19	AAW47623
4	117	79.1	28	19	AAW47623
5	117	79.1	28	19	AAW47623
6	117	79.1	28	20	AAW47623
7	117	79.1	28	20	AAW47623
8	117	79.1	28	20	AAW47623
9	112	75.7	28	19	AAW47614
10	112	75.7	28	20	AAW47614
11	112	75.7	338	22	AAU14026

12	112	75.7	345	21	AAW4536	HIV-1 isolate LAI
13	112	75.7	345	22	AAW4536	Amino acid sequenc
14	112	75.7	420	15	AAW43785	Translation of HIV
15	112	75.7	853	19	AAW43066	HIV-1 gp120 protei
16	112	75.7	856	14	AAW41025	Selectively deglyc
17	112	75.7	856	14	AAW41026	Selectively deglyc
18	112	75.7	856	14	AAW41027	Selectively deglyc
19	112	75.7	856	14	AAW41028	Selectively deglyc
20	112	75.7	856	14	AAW41029	Selectively deglyc
21	112	75.7	856	14	AAW41030	Selectively deglyc
22	112	75.7	856	14	AAW41031	Selectively deglyc
23	112	75.7	856	14	AAW41032	Selectively deglyc
24	112	75.7	856	21	AAW47072	Wild type HIV-1 HA
25	112	75.7	856	22	AAW45697	HIV-1/11B env clo
26	112	75.7	856	19	AAW43067	HIV-1 gp120 protei
27	112	75.7	863	13	AAW48955	Non-cleavable, sol
28	112	75.7	865	16	AAW43909	HIV-1 envelope pol
29	112	75.7	868	7	AAW60063	HIV virus env gene
30	112	75.7	868	7	AAW60422	Sequence of LAV vi
31	112	75.7	901	8	AAW7065	Sequence encoded b
32	108	73.0	28	19	AAW47624	Antimicrobial pept
33	108	73.0	28	19	AAW47634	Antimicrobial pept
34	108	73.0	28	20	AAW47635	Antimicrobial pept
35	108	73.0	28	20	AAW47635	Antimicrobial pept
36	107	72.3	28	19	AAW47625	Antimicrobial pept
37	107	72.3	28	19	AAW47771	Antimicrobial pept
38	107	72.3	28	19	AAW47772	Antimicrobial pept
39	107	72.3	28	19	AAW47635	Antimicrobial pept
40	107	72.3	28	20	AAW432705	Antimicrobial pept
41	107	72.3	28	20	AAW432706	Antimicrobial pept
42	107	72.3	28	20	AAW432561	Antimicrobial pept
43	107	72.3	412	11	AAW5095	Antimicrobial pept
44	107	72.3	412	11	AAW5095	Synthetic HIV-1 tr
45	107	72.3	704	11	AAW5096	PSD302.PEP HIV-1 g

ALIGNMENTS

RESULT	1
AAW47769	standard; peptide: 28 AA.
ID	AAW47769; standard; peptide: 28 AA.
AC	AAW47769;
XX	
DT	26-MAY-1998 (first entry)
XX	
DE	Antimicrobial peptide LLPI analogue.
XX	
KW	Antimicrobial; transmembrane protein; TM; lentivirus lytic peptide;
KW	LLPI; amphipathic; antibacterial; antifungal; antiviral; antiprotzoal.
XX	
OS	Synthetic.
OS	Human immunodeficiency virus.
XX	
PN	US5714577-A.
XX	
PD	03-FEB-1998.
XX	
PF	24-JAN-1997; 97US-0786748.
XX	
PR	26-JAN-1996; 96US-0010634.
XX	
PA	24-JAN-1997; 97US-0786748.
XX	
PI	(UYP1-) UNIV PITTSBURGH.
XX	
PI	Mietzner TA, Montelaro RC, Tencza SB;
XX	
DR	WPI; 1998-158352/14.
XX	
PT	Retroviral TM peptides - useful as antibacterial agents
XX	
PS	Disclosure; Column 19; 59pp; English.

XX The invention relates to new antimicrobial peptides which correspond to
CC amino acid sequences in the transmembrane proteins of lentiviruses, in
CC particular HIV and SIV. These peptides comprise arginine rich sequences
CC which, when modelled for secondary structure, display high
CC amphipathicity and hydrophobic moment. Also disclosed are structural
CC and functional analogues and homologues of these peptides which also
CC display antimicrobial activity. The peptides are highly inhibitory to
CC microorganisms (bacteria, fungi, viruses and protozoa) but significantly
CC less toxic to red blood cells and other normal mammalian cells. Activity
CC is demonstrated against Gram positive and negative bacteria including
CC *Pseudomonas aeruginosa*, *Staphylococcus aureus*, *Enterococcus faecalis* and
CC *Serratia marcescens*.
CC The present sequence is one of 169 disclosed specific examples of
CC the new peptides. It is an analogue of the peptide designated LLPI
CC (see AAM47614) which is a peptide from the transmembrane protein (gp41)
CC of HIV strain HXB2R.
XX
SQ Sequence 28 AA:

Query Match 82.4%; Score 122; DB 19; Length 28;
Best Local Similarity 92.9%; Pred. No. 1.5e-10;
Matches 26; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 1 RVIRVQACRAIRHIVRIRIGLRRL 28
1 RIVRVGACRAIRHIVRIRIGLRRL 28
DB 1 RIVRVGACRAIRHIVRIRIGLRRL 28

RESULT 2
AAY32703
ID AAY32703 standard; peptide: 28 AA.
XX
AC AAY32703:
XX
DT 21-OCT-1999 (first entry)
XX
DE Antimicrobial peptide LLPI analogue.
XX
KW Antimicrobial peptide: LLPI; SLP-1; LPP2; SLP2A; SLP2B; ELP; Infection;
KW growth inhibitor; microorganism; virus; gene therapy; vector production;
KW sterilisation.
XX
OS Synthetic.
OS Human immunodeficiency virus type 1.
XX
PN US5945507-A.
XX
PD 31-AUG-1999.
XX
PF 18-SEP-1997; 97US-0932682.
XX
PR 26-JAN-1996; 96US-0010634.
PR 24-JAN-1997; 97US-0786748.
PR 18-SEP-1997; 97US-0932682.
XX
PA (UVP1-) UNIV PITTSBURGH.
XX
PI Metzner TA, Montelaro RC, Tencza SB;
XX
DR WPI; 1999-508189/42.
XX
PT Antimicrobial peptides useful for treating microbial infections
XX
PS Disclosure; Column 21; 62pp; English.
XX
CC This sequence represents an antimicrobial peptide of the invention, and
CC is an analogue of the peptide LLPI (see AAY32549). The peptides can be
CC used for treating infections caused by *Staphylococcus aureus*,
CC microorganisms (bacteria, fungi, viruses and protozoa) but significantly
CC less toxic to red blood cells and other normal mammalian cells. Activity
CC is demonstrated against Gram positive and negative bacteria including
CC *Pseudomonas aeruginosa*, *Staphylococcus aureus*, *Enterococcus faecalis*
CC and *Serratia marcescens*, *Escherichia coli*, fungi, protozoa and viruses in
CC a mammalian host. They can be used to inhibit growth of diverse
CC

CC microorganisms such as bacteria, fungi, protozoa and DNA and RNA viruses
CC and can be used in tissue culture to inhibit unwanted microbial growth,
CC particularly for the production of recombinant proteins or vectors for
CC gene therapy. They can also be used in preventing infections through the
CC sterilisation of wounds prior to suture and to sterilise surgical
CC instruments. The unique structure of these antimicrobial peptides
CC imparts high potency while selectivity is maintained. They are
CC moderately haemolytic but only lyse red blood cells at high
CC concentrations unlike melittin, a peptide extracted from bee venom, which
CC is highly active against bacteria and lyses red blood cells showing
CC little selectivity. The peptides target a membrane structure which makes
CC it more difficult for a microorganism to develop a mechanism of
CC resistance against this type of antibiotic. Their small size makes them
CC relatively simple to prepare by standard synthetic peptide chemistry.
XX
SQ Sequence 28 AA:

Query Match 82.4%; Score 122; DB 20; Length 28;
Best Local Similarity 92.9%; Pred. No. 1.5e-10;
Matches 26; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 1 RVIRVQACRAIRHIVRIRIGLRRL 28
1 RIVRVGACRAIRHIVRIRIGLRRL 28
DB 1 RIVRVGACRAIRHIVRIRIGLRRL 28

RESULT 3
AAM47623
ID AAM47623 standard; peptide: 28 AA.
XX
AC AAM47623:
XX
DT 26-MAY-1998 (first entry)
XX
DE Antimicrobial peptide LLPI analogue.
XX
KW Antimicrobial; transmembrane protein; TM; lentivirus lytic peptide;
KW LPI; amphipathic; antibacterial; antifungal; antiviral; antiprotozoal.
XX
OS Synthetic.
OS Human immunodeficiency virus.
XX
PN US5714577-A.
XX
PD 03-FEB-1998.
XX
PF 24-JAN-1997; 97US-0786748.
XX
PR 26-JAN-1996; 96US-0010634.
PR 24-JAN-1997; 97US-0786748.
XX
PA (UVP1-) UNIV PITTSBURGH.
XX
PI Metzner TA, Montelaro RC, Tencza SB;
XX
DR WPI; 1998-158352/14.
XX
PT Retroviral TM peptides - useful as antibacterial agents
XX
PS Disclosure; Column 9; 59pp; English.
XX
CC The invention relates to new antimicrobial peptides which correspond to
CC amino acid sequences in the transmembrane proteins of lentiviruses, in
CC particular HIV and SIV. These peptides comprise arginine rich sequences
CC which, when modelled for secondary structure, display high
CC amphipathicity and hydrophobic moment. Also disclosed are structural
CC and functional analogues and homologues of these peptides which also
CC display antimicrobial activity. The peptides are highly inhibitory to
CC microorganisms (bacteria, fungi, viruses and protozoa) but significantly
CC less toxic to red blood cells and other normal mammalian cells. Activity
CC is demonstrated against Gram positive and negative bacteria including
CC *Pseudomonas aeruginosa*, *Staphylococcus aureus*, *Enterococcus faecalis* and

CC Serratia marcescens.
CC The present sequence is one of 169 disclosed specific examples of
CC the new peptides. It is an analogue of the peptide designated LLP1
CC (see AAW47614) which is a peptide from the transmembrane protein (gp41)
CC of HIV strain HXB2R.
XX
SQ Sequence 28 AA;

Query Match 79.1%; Score 117; DB 19; Length 28;
Best Local Similarity 89.3%; Pred. No. 7.8e-10;
Matches 25; Conservative 0; Mismatches 3; Indels 0; Gaps 0;
QY 1 RVIRVVQACRAIRHIVRRIRGRLRL 28
DB 1 rvirvvgacrairhivrrirglerll 28

RESULT 4
AAW47628
ID AAW47628 standard; peptide; 28 AA.
XX
AC AAW47628;
XX
DT 26-MAY-1998 (first entry)
XX
DE Antimicrobial peptide LLP1 analogue.
XX
KW Antimicrobial; transmembrane protein; TM; lentivirus lytic peptide;
XX LLP; amphipathic; antibacterial; antifungal; antiviral; antiprotozoal.
OS Synthetic.
XX
OS Human immunodeficiency virus.
XX
PN US5714577-A.
XX
PD 03-FEB-1998.
XX
PF 24-JAN-1997; 97US-0786748.
XX
PR 26-JAN-1996; 96US-0010634.
XX
PR 24-JAN-1997; 97US-0786748.
XX
PA (UYPI-) UNIV PITTSBURGH.
PI Mietzner TA, Montelaro RC, Tencza SB;
XX
DR WPI; 1998-158352/14.
XX
PT Retroviral TM peptides - useful as antibacterial agents
XX
PS Disclosure; Column 9; 59pp; English.
XX
CC The invention relates to new antimicrobial peptides which correspond to
CC amino acid sequences in the transmembrane proteins of lentiviruses, in
CC particular HIV and SIV. These peptides comprise arginine rich sequences
CC which, when modelled for secondary structure, display high
CC amphipathicity and hydrophobic moment. Also disclosed are structural
CC and functional analogues and homologues of these peptides which also
CC display antimicrobial activity. The peptides are highly inhibitory to
CC microorganisms (bacteria, fungi, viruses and protozoa) but significantly
CC less toxic to red blood cells and other normal mammalian cells. Activity
CC is demonstrated against Gram positive and negative bacteria including
CC pseudomonas aeruginosa, Staphylococcus aureus, Enterococcus faecalis and
CC Serratia marcescens.
CC The present sequence is one of 169 disclosed specific examples of
CC the new peptides. It is an analogue of the peptide designated LLP1
CC (see AAW47614) which is a peptide from the transmembrane protein (gp41)
CC of HIV strain HXB2R.
XX
SQ Sequence 28 AA;

Query Match 79.1%; Score 117; DB 19; Length 28;
Best Local Similarity 89.3%; Pred. No. 7.8e-10;
Matches 25; Conservative 0; Mismatches 3; Indels 0; Gaps 0;
QY 1 RVIRVVQACRAIRHIVRRIRGRLRL 28
DB 1 rvirvvgacrairhivrrirglerll 28

RESULT 5
AAW47633
ID AAW47633 standard; peptide; 28 AA.
XX
AC AAW47633;
XX
DT 26-MAY-1998 (first entry)
XX
DE Antimicrobial peptide LLP1 analogue.
XX
KW Antimicrobial; transmembrane protein; TM; lentivirus lytic peptide;
XX LLP; amphipathic; antibacterial; antifungal; antiviral; antiprotozoal.
OS Synthetic.
XX
OS Human immunodeficiency virus.
XX
PN US5714577-A.
XX
PD 03-FEB-1998.
XX
PF 24-JAN-1997; 97US-0786748.
XX
PR 26-JAN-1996; 96US-0010634.
XX
PR 24-JAN-1997; 97US-0786748.
XX
PA (UYPI-) UNIV PITTSBURGH.
PI Mietzner TA, Montelaro RC, Tencza SB;
XX
DR WPI; 1998-158352/14.
XX
PT Retroviral TM peptides - useful as antibacterial agents
XX
PS Disclosure; Column 9; 59pp; English.
XX
CC The invention relates to new antimicrobial peptides which correspond to
CC amino acid sequences in the transmembrane proteins of lentiviruses, in
CC particular HIV and SIV. These peptides comprise arginine rich sequences
CC which, when modelled for secondary structure, display high
CC amphipathicity and hydrophobic moment. Also disclosed are structural
CC and functional analogues and homologues of these peptides which also
CC display antimicrobial activity. The peptides are highly inhibitory to
CC microorganisms (bacteria, fungi, viruses and protozoa) but significantly
CC less toxic to red blood cells and other normal mammalian cells. Activity
CC is demonstrated against Gram positive and negative bacteria including
CC pseudomonas aeruginosa, Staphylococcus aureus, Enterococcus faecalis and
CC Serratia marcescens.
CC The present sequence is one of 169 disclosed specific examples of
CC the new peptides. It is an analogue of the peptide designated LLP1
CC (see AAW47614) which is a peptide from the transmembrane protein (gp41)
CC of HIV strain HXB2R.
XX
SQ Sequence 28 AA;

Query Match 79.1%; Score 117; DB 19; Length 28;
Best Local Similarity 89.3%; Pred. No. 7.8e-10;
Matches 25; Conservative 0; Mismatches 3; Indels 0; Gaps 0;
QY 1 RVIRVVQACRAIRHIVRRIRGRLRL 28
DB 1 rvirvvgacrairhivrrirglerll 28

xx	AAI32559;
xx	21-OCT-1999 (first entry)
xx	Antimicrobial peptide LLPI analogue.
xx	Antimicrobial peptide; LLPI; SLP-1; LLP2; SUP2A; SUP2B; ELP; infection;
xx	growth inhibitor; microorganism; virus; gene therapy; vector production;
xx	sterilisation.
xx	Synthetic.
xx	Human immunodeficiency virus type 1.
xx	US5945507-A.
xx	31-AUG-1999.
xx	18-SEP-1997; 97US-0932682.
xx	26-JAN-1996; 96US-0010634.
xx	24-JAN-1997; 97US-0786748.
xx	18-SEP-1997; 97US-0932682.
xx	(UYP1-) UNIV PITTSBURGH.
xx	Mietzner TA, Montelaro RC, Tencza SB;
xx	WPI: 1999-508189/42.
xx	Antimicrobial peptides useful for treating microbial infections
xx	Disclosure: Column 9; 62pp; English.
xx	This sequence represents an antimicrobial peptide of the invention, and
xx	is an analogue of the peptide LLPI (see AAI32549). The peptides can be
xx	used for treating infections caused by <i>Staphylococcus aureus</i> ,
xx	methicillin resistant <i>S. aureus</i> , <i>Pseudomonas aeruginosa</i> , <i>Enterococcus</i>
xx	<i>faecalis</i> , <i>S. marescens</i> , <i>Escherichia coli</i> , fungi, protozoa and viruses in
xx	a mammalian host. They can be used to inhibit growth of diverse
xx	microorganisms such as bacteria, fungi, protozoa and DNA and RNA viruses
xx	and can be used in tissue culture to inhibit unwanted microbial growth,
xx	particularly for the production of recombinant proteins or vectors for
xx	gene therapy. They can also be used in preventing infections through the
xx	sterilisation of wounds prior to suture and to sterilise surgical
xx	instruments. The unique structure of these antimicrobial peptides
xx	imparts high potency while selectivity is maintained, they are
xx	moderately haemolytic but only lyse red blood cells at high
xx	concentrations unlike melittin, a peptide extracted from bee venom, which
xx	is highly active against bacteria and lyses red blood cells showing
xx	little selectivity. The peptides target a membrane structure which makes
xx	it more difficult for a microorganism to develop a mechanism of
xx	resistance against this type of antibiotic. Their small size makes them
xx	relatively simple to prepare by standard synthetic peptide chemistry.
xx	Sequence 28 AA:
OY	Query Match 79.1%; Score 117; DB 20; Length 28;
Db	Best Local Similarity 89.3%; Pred. No. 7.8e-10;
	Matches 25; Conservative 0; Mismatches 3; Indels 0; Gaps 0
	1 RVIRVQRCRAIRHVRIRGRLRIL 28
	1 rvlrvvgqacrairhprlrgrlerll 28

XX		AAI32564;	
AC		21-OCT-1999	(first entry)
DT		Antimicrobial peptide LLP1 analogue.	
DE		Antimicrobial peptide LLP1 analogue.	
XX		Antimicrobial peptide; LLP1; SLP-1; LLP2; SUP2A; SLP2B; ELP; infection; growth inhibitor; microorganism; virus; gene therapy; vector production; sterilisation.	
KM		Synthetic.	
OS		Human immunodeficiency virus type 1.	
XX		US5945507-A.	
PN		31-AUG-1999.	
PD		18-SEP-1997;	97US-0932682.
XX		26-JAN-1996;	96US-0010634.
PR		24-JAN-1997;	97US-0786748.
PR		18-SEP-1997;	97US-0932682.
XX		(UYP1-) UNIV PITTSBURGH.	
PA		Meltzer TA,	Montelaro RC,
PI		Tencza SB;	
XX		WPI; 1999-508189/42.	
DR		Antimicrobial peptides useful for treating microbial infections	
XX		Disclosure; Column 9; 62pp: English.	
PS		This sequence represents an antimicrobial peptide of the invention, and is an analogue of the peptide LLP1 (see AAI32564). The peptides can be used for treating infections caused by <i>Staphylococcus aureus</i> , <i>Methicillin resistant S. aureus</i> , <i>Pseudomonas aeruginosa</i> , <i>Enterococcus faecalis</i> , <i>S. marcescens</i> , <i>Escherichia coli</i> , fungi, protozoa and viruses in a mammalian host. They can be used to inhibit growth of diverse microorganisms such as bacteria, fungi, protozoa and DNA and RNA viruses and can be used in tissue culture to inhibit unwanted microbial growth, particularly for the production of recombinant proteins or vectors for gene therapy. They can also be used in preventing infections through the sterilisation of wounds prior to suture and to sterilise surgical instruments. The unique structure of these antimicrobial peptides imparts high potency while selectivity is maintained, they are moderately haemolytic but only lyse red blood cells at high concentrations unlike melittin, a peptide extracted from bee venom, which is highly active against bacteria and lyses red blood cells showing little selectivity. The peptides target a membrane structure which makes it more difficult for a microorganism to develop a mechanism of resistance against this type of antibiotic. Their small size makes them relatively simple to prepare by standard synthetic peptide chemistry.	
CC		Sequence	28 AA;
SQ			
Query Match		79.1%;	Score 117; DB 20;
Best Local Similarity		89.3%;	Pred No. 7.8e-10;
Matches	25;	Conservative	0;
		Mismatches	3;
		Indels	0;
		Gaps	0;
OY	1	RVIKVVORACRAIRHIVRRINGRLRLL 28	
Db	1	rviivvgacrairnhprirgrrll 28	
RESULT	8		
AAI32569			
ID	AAI32569	standard; peptide: 28 AA.	
AC	AAI32569;		
XX			
XX			

```

DT 21-OCT-1999 (first entry)
XX Antimicrobial peptide LRP1 analogue.
DE Antimicrobial peptide: LRP1; SLP-1; LRP2; SLP2A; SLP2B; ELP; infection;
XX growth inhibitor; microorganism; virus; gene therapy; vector production;
KM sterilisation.
XX Synthetic.
OS Human immunodeficiency virus type 1.
XX US5945507-A.
XX 31-AUG-1999.
XX 18-SEP-1997; 97US-0932682.
XX 26-JAN-1996; 96US-0010634.
PR 24-JAN-1997; 97US-0786748.
PR 18-SEP-1997; 97US-0932682.
XX (UYP1-) UNIV PITTSBURGH.
XX Mletzner TA, Montelaro RC, Tencza SB;
XX WPI; 1999-508189/42.
XX Antimicrobial peptides useful for treating microbial infections
XX Disclosure; Column 9; 62pp; English.
XX This sequence represents an antimicrobial peptide of the invention, and
XX is an analogue of the peptide LRP1 (see AAY32549). The peptides can be
XX used for treating infections caused by staphylococcus aureus,
XX methicillin resistant S. aureus, Pseudomonas aeruginosa, Enterococcus
XX faecalis, S. marcescens, Escherichia coli, fungi, protozoa and viruses in
XX a mammalian host. They can be used to inhibit growth of diverse
XX microorganisms such as bacteria, fungi, protozoa and DNA and RNA viruses
XX and can be used in tissue culture to inhibit unwanted microbial growth,
XX particularly for the production of recombinant proteins or vectors for
XX gene therapy. They can also be used in preventing infections through the
XX sterilisation of wounds prior to suture and to sterilise surgical
XX instruments. The unique structure of these antimicrobial peptides
XX imparts high potency while selectivity is maintained, they are
XX moderately haemolytic but only lyse red blood cells at high
XX concentrations unlike melittin, a peptide extracted from bee venom, which
XX is highly active against bacteria and lyses red blood cells showing
XX little selectivity. The peptides target a membrane structure which makes
XX it more difficult for a microorganism to develop a mechanism of
XX resistance against this type of antibiotic. Their small size makes them
XX relatively simple to prepare by standard synthetic peptide chemistry.
XX Sequence 28 AA;
SQ

```

```

Query Match 79.1%; Score 117; DB 20; Length 28;
Best Local Similarity 89.3%; Pred. No. 7.8e-10;
Matches 25; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

```

```

OY 1 RVIRVORACRAIRHYRIRROGLRRL 28
    ||||| ||||| ||||| ||||| |||
DB 1 RVIRVVGACRAIRHPRIIRIGLERIL 28

```

```

RESULT 9
ID AAY47614
AAW47614 standard; peptide; 28 AA.
XX AAY47614;
XX AC
XX 26-MAY-1998 (first entry)
XX DT
XX Antimicrobial peptide HIVXB2R 828-855, or LRP1.
DE

```

```

XX Antimicrobial; transmembrane protein; TM; lentivirus lytic peptide;
KM LRP; amphipathic; antibacterial; antifungal; antiviral; antiprotzoal.
XX Human immunodeficiency virus.
XX US5714577-A.
XX 03-FEB-1998.
XX 24-JAN-1997; 97US-0786748.
XX 26-JAN-1996; 96US-0010634.
PR 24-JAN-1997; 97US-0786748.
XX (UYP1-) UNIV PITTSBURGH.
XX Mletzner TA, Montelaro RC, Tencza SB;
XX WPI; 1998-158352/14.
XX Retroviral TM peptides - useful as antibacterial agents
XX Disclosure; Column 5; 59pp; English.
XX The invention relates to new antimicrobial peptides which correspond to
XX amino acid sequences in the transmembrane proteins of lentiviruses, in
XX particular HIV and SIV. These peptides comprise arginine rich sequences
XX which, when modelled for secondary structure, display high
XX amphipathicity and hydrophobic moment. Also disclosed are structural
XX and functional analogues and homologues of these peptides which also
XX display antimicrobial activity. The peptides are highly inhibitory to
XX microorganisms (bacteria, fungi, viruses and protozoa) but significantly
XX less toxic to red blood cells and other normal mammalian cells. Activity
XX is demonstrated against gram positive and negative bacteria including
XX Pseudomonas aeruginosa, Staphylococcus aureus, Enterococcus faecalis and
XX Serratia marcescens.
XX The present sequence is one of 169 disclosed specific examples of
XX the new peptides. It is called LRP1 and corresponds to residues 828-855
XX of the transmembrane protein (gp41) of HIV strain HXB2R.
XX Sequence 28 AA;
SQ

```

```

Query Match 75.7%; Score 112; DB 19; Length 28;
Best Local Similarity 85.7%; Pred. No. 4e-09;
Matches 24; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

```

```

OY 1 RVIRVORACRAIRHYRIRROGLRRL 28
    ||| ||| ||||| ||||| ||||| |||
DB 1 RVIRVVGACRAIRHPRIIRIGLERIL 28

```

```

RESULT 10
ID AAY32549
AAY32549 standard; peptide; 28 AA.
XX AAY32549;
XX AC
XX 21-OCT-1999 (first entry)
XX DT
XX Antimicrobial peptide LRP1.
DE

```

```

XX Antimicrobial peptide: LRP1; SLP-1; LRP2; SLP2A; SLP2B; ELP; infection;
KM growth inhibitor; microorganism; virus; gene therapy; vector production;
KM sterilisation.
XX Human immunodeficiency virus type 1.
XX OS
XX US5945507-A.
XX PN
XX 31-AUG-1999.
XX PD
XX

```

PF	18-SEP-1997;	97US-0932682.	
XX			
PR	26-JAN-1996;	96US-0010634.	
PR	24-JAN-1997;	97US-0786748.	
PR	18-SEP-1997;	97US-0932682.	
XX			
PA	(UYPI-) UNIV PITTSBURGH.		
XX			
P1	Mietzner TA, Montelaro RC, Tenenza SB;		
XX			
DR	WPI: 1999-508189/42.		
XX			
PT	Antimicrobial peptides useful for treating microbial infections		
XX			
PS	Example 1; Column 5; 62pp; English.		
XX			
CC	This sequence represents the antimicrobial peptide LRP1, and was used		
CC	to design the peptide analogues of the invention. The peptides can be		
CC	used for treating infections caused by <i>Staphylococcus aureus</i> , <i>methicillin</i>		
CC	resistant <i>S. aureus</i> , <i>Pseudomonas aeruginosa</i> , <i>Enterococcus faecalis</i> ,		
CC	<i>S. marcescens</i> , <i>Escherichia coli</i> , fungi, protozoa and viruses in a		
CC	mammalian host. They can be used to inhibit growth of diverse		
CC	microorganisms such as bacteria, fungi, protozoa and DNA and RNA viruses		
CC	and can be used in tissue culture to inhibit unwanted microbial growth,		
CC	particularly for the production of recombinant proteins or vectors for		
CC	gene therapy. They can also be used in preventing infections through the		
CC	sterilisation of wounds prior to suture and to sterilise surgical		
CC	instruments. The unique structure of these antimicrobial peptides imparts		
CC	high potency while selectivity is maintained, they are moderately		
CC	haemolytic but only lyse red blood cells at high concentrations unlike		
CC	melittin, a peptide extracted from bee venom, which is highly active		
CC	against bacteria and lyses red blood cells showing little selectivity.		
CC	The peptides target a membrane structure which makes it more difficult		
CC	for a microorganism to develop a mechanism of resistance against this		
CC	type of antibiotic. Their small size makes them relatively simple to		
CC	prepare by standard synthetic peptide chemistry.		
XX			
SO	Sequence 28 AA:		
	Query Match 75.7%; Score 112; DB 20; Length 28;		
	Best Local Similarity 85.7%; Pred No. 4e-09;		
	Matches 24; Conservative 0; Mismatches 4; Indels 0; Gaps 0		
QY	1 RYIRVVQRACRAIRHIVRRIRROGLRRITL 28		
Db	1 rlievvgagcainrhprirgrlerll 28		
	RESULT 11		
ID	AAU14026		
XX	AAU14026 standard: peptide; 338 AA.		
XX			
AC	AAU14026;		
XX			
DT	21-NOV-2001 (first entry)		
XX			
DE	Peptide sequence from HIV-1 isolate BRU enveloped protein gp41.		
XX			
KW	Anti-retroviral; DP178-like; DP107-like; transmembrane protein gp41;		
RW	antifusogenic; antiviral; HIV transmission.		
XX			
OS	Human immunodeficiency virus type 1 isolate BRU.		
XX			
PN	WO200151673-A2.		
PD	19-JUL-2001.		
XX			
PF	05-JUL-2000; 2000WO-US35727.		
XX			
RR	09-JUL-1999; 99US-0350841.		
XX			
PA	(TRIM-) TRIMERIS INC.		

XX	Jeffs P, Lackey JW, Erickson JB, Lawless MK, Merutka G;
PI	WP1: 2001-442157/47.
DR	
XX	
PT	Identifying a compound that inhibits the formation of or disrupts a
PR	Dp107/Dp178 complex, especially compounds with antifeugenic, antiviral
PT	or intracellular modulatory activity, by detecting the formation of a
PT	Dp107/Dp178 complex -
XX	
PS	Disclosure: Fig 20; 259pp; English.
XX	
CC	The present invention relates to peptides which exhibit anti-retroviral
CC	activity. The peptides of the invention (AAU12559-AAU14009) comprise
CC	Dp178-like and Dp107-like peptides. The Dp178 peptide corresponds
CC	to amino acids 639-673 of the transmembrane protein gp41 from human
CC	immunodeficiency virus 1 (HIV-1) isolate LAI. The Dp107 peptide
CC	corresponds to amino acids 558-595 of gp41 from HIV-1LAI. The invention
CC	also relates to a method of identifying compounds that inhibit the
CC	formation of or disrupts a Dp107/Dp178 complex. The method comprises
CC	detecting the formation of a Dp107/Dp178 complex, both in the presence
CC	or absence of a test compound, in a reaction mixture containing Dp107
CC	and Dp178 peptides. The method is useful for identifying compounds,
CC	including small molecule compounds, which may themselves exhibit
CC	antifeugenic, antiviral or intracellular modulatory activity. The
CC	Dp178-like/Dp107-like peptides are useful to inhibit human and non-human
CC	retroviral, particularly HIV, transmission to uninfected cells. The
CC	present sequence represents a peptide sequence from HIV-1 isolate
CC	BRU envelope protein gp41.
XX	
SQ	Sequence 338 AA;
	Query Match 75.7%; Score 112; DB 22; Length 338;
	Best Local Similarity 85.7%; Pred. No. 5e-08;
	Matches 24; Conservative 0; Mismatches 4; Indels 0; Gaps 0;
OY	1 RVIKRVQRCRAIRHIVRIRIGLRRL 28
DB	310 rlvievvgacrairhiviririglerl 337
RESULT 12	
AA014536	
ID	AA014536 standard; Protein; 345 AA.
XX	
AC	AA014536;
XX	
DT	24-NOV-2000 (first entry)
XX	
DE	HIV-1 isolate LAI gp41 protein.
XX	
KW	HIV-1; gp41; N-helical domain; heptad repeat region; C-helical domain;
KW	gp41 transmembrane-proximal amphipathic alpha-helical segment;
KW	core 6-helix bundle; viral entry inhibition; immunogenic;
KW	antibody: humoral response; broad spectrum vaccine; anti-HIV;
KW	envelope glycoprotein; prophylaxis; therapy; group M; subtype B;
KW	isolate LAI.
XX	
OS	Human immunodeficiency virus type 1.
XX	
PN	WO200040616-A1.
PD	
PD	13-JUL-2000.
XX	
PF	10-JAN-2000; 2000WO-US00456.
XX	
PR	08-JAN-1999; 99US-0115404.
PR	07-JAN-2000; 2000US-0480336.
XX	
PA	(WILD/) WILD C T.
PA	(WEIS/) WEISS C D.
XX	

XX (HARD) HARVARD COLLEGE.
 PA Essex ME, Lee TH, Yu X;
 PI WPI; 1994-200197/24.
 DR N-PSDB; AA066275.
 XX
 PT Method for treating HIV patients - comprises administration of
 mutated gp41 polypeptide
 XX
 PS Claim 11; Fig 1; 54pp; English.
 CC The inventors claim a method of treating a patient infected with HIV
 CC by administering a mutated gp41 polypeptide or a therapeutic
 CC composition comprising nucleic acid encoding the mutant gp41
 CC polypeptide in an expressible genetic construction. The mutant gp41
 CC of the following regions of wild type gp41 (AAR53783): AAs 844-856;
 CC 814-856; 796-856; 776-856; 753-856; or 710-856, effective to either
 CC disrupt viral replication or HIV or disrupt the assembly of viral
 CC Env proteins in an HIV infected cell. AA066275-corresp. to bps 7631-
 CC 8890 of wt HIV-1. X in the AA sequence represents the posn. of a
 CC stop codon in AA066275.
 CC
 SQ Sequence 420 AA;

Query Match 75.7%; Score 112; DB 15; Length 420;
 Best Local Similarity 85.7%; Pred. No. 6.2e-08;
 Matches 24; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

OY 1 RVIRVORACRAIRHIVRIROGLRLIL 28
 ||| ||| ||||| ||||| |||
 Db 359 RVLEVVGACRAIRHPIRIRGLRLIL 386

RESULT 15

AAW43066
 ID AAW43066 standard; peptide; 853 AA.

AC AAW43066;

DT 11-SEP-1998 (first entry)

DE HIV-1 gp120 protein fragment from isolate HXB2.

KW gp120 protein; purification; fractionation; ion exchange; chromatography;
 binding affinity; CD4; hydrophobic interaction; size exclusion; vaccine.

OS Human immunodeficiency virus type 1.

PN US5696238-A.

PD 09-DEC-1997.

PF 11-MAY-1995; 95US-0439286.

PR 20-AUG-1991; 91US-0684963.

PR 16-AUG-1993; 93US-0109002.

PR 09-MAY-1994; 94US-0240073.

PR 11-MAY-1995; 95US-0439286.

PA (CHIR) CHIRON CORP.

PI Haigwood NL, Scandella C;

DR WPI; 1998-041353/04.

PT Purification of HIV gp120 - using chromatographic methods
 PS Disclosure; Fig 2A-W; 53pp; English.
 XX

CC AAW43066-M43080 are fragments of the gp120 protein from different human
 CC immunodeficiency virus type I (HIV-1) isolates. These proteins are used
 CC in a novel method for purifying HIV gp120 so as to provide a purified
 CC gp120 glycoprotein having protein/protein binding properties
 CC substantially identical to natural viral HIV gp120. The method involves
 CC fractionating a crude gp120 preparation containing full-length,
 CC glycosylated gp120 using ion exchange chromatography so as to provide a
 CC first collection of fractions. A fraction from the first collection is
 CC selected that exhibits specific binding affinity for CD4 peptide,
 CC thereby producing a first fractionated material. The first fractionated
 CC material is fractionated by hydrophobic interaction chromatography so as
 CC to provide a second collection of fractions from which a second
 CC collection is selected that exhibits specific binding affinity for CD4
 CC peptide. This second fraction is fractionated by size exclusion
 CC chromatography so as to provide a third collection of fractions
 CC exhibiting specific binding affinity for CD4 peptide, thereby providing
 CC the purified gp120. The purified gp120 can be used for antibody
 CC production and in vaccines.
 CC
 SQ Sequence 853 AA;

Query Match 75.7%; Score 112; DB 19; Length 853;
 Best Local Similarity 85.7%; Pred. No. 1.3e-07;
 Matches 24; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

OY 1 RVIRVORACRAIRHIVRIROGLRLIL 28
 ||| ||| ||||| ||||| |||
 Db 825 RVLEVVGACRAIRHPIRIRGLRLIL 852

Search completed: August 14, 2002, 10:50:38
 Job time: 344 sec